STIC Search Report
Biotech-Chem Library

STIC Database Tracking Number: 168058

TO:	From:
Celine Qian **Location: rem/2A64/2C70** **Art Unit: 1636** **Wednesday, October 19, 2005** **Case Serial Number: 09/877935**	**Edward Hart** **Location: Biotech-Chem Library** **REM-1A55** **Phone: 571-272-2512** **edward.hart@uspto.gov**

Search Notes

Examiner Qian,

Here are the results of the search you requested.

Please feel free to contact me if you have any questions.

Edward Hart

ATTN: Ed Hurt

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CRFE

Scientific and Technical Information Center

SEARCH REQUEST FORM

Requester's Full Name: Celine Qian Examiner #: 78770 Date: 10/14/05

Art Unit: 1636 Phone Number: 2-0777 Serial Number: 09/877935

Location (Bldg/Room#): 2164 (Mailbox #): 2C70 Results Format Preferred (circle): ~~PAPER~~ DISK

MEY

To ensure an efficient and quality search, please attach a copy of the cover sheet, claims, and abstract or fill out the following:

Title of Invention: Regulatory sequences of the mouse villin gene

Inventors (please provide full names): Pinto et al.

Earliest Priority Date: 10/9/1998

Search Topic:

Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc., if known.

For Sequence Searches Only *Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number.*

Please search SEQ ID NO: 1, (interference only.)

NA 8995

MEY

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STIC

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****

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Searcher: _____

Searcher Phone #: _____

Searcher Location: _____

Date Searcher Picked Up: 10/18/05

Date Completed: _____

Searcher Prep & Review Time: _____

Online Time: _____

Type of Search

1 NA Sequence (#)

___ AA Sequence (#)

___ Structure (#)

___ Bibliographic

___ Litigation

___ Fulltext

___ Other

Vendors and cost where applicable

___ STN ___ Dialog

___ Questel/Orbit ___ Lexis/Nexis

___ Westlaw ___ WWW/Internet

104 In-house sequence systems

___ Commercial ___ Oligomer ___ Score/Length

X Interference ___ SPDI ___ Encode/Translate

___ Other (specify)

STIC SEARCH RESULTS FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact *the searcher or contact*:

Mary Hale, Information Branch Supervisor
Remsen Bldg. 01 D86
571-272-2507

Voluntary Results Feedback Form

- *I am an examiner in Workgroup:* ☐ *Example: 1610*
- *Relevant prior art **found**, search results used as follows:*
  - ☐ 102 rejection
  - ☐ 103 rejection
  - ☐ Cited as being of interest.
  - ☐ Helped examiner better understand the invention.
  - ☐ Helped examiner better understand the state of the art in their technology.

Types of relevant prior art found:

- ☐ Foreign Patent(s)
- ☐ Non-Patent Literature
  (journal articles, conference proceedings, new product announcements etc.)

- *Relevant prior art **not found**:*
  - ☐ Results verified the lack of relevant prior art (helped determine patentability).
  - ☐ Results were not useful in determining patentability or understanding the invention.

Comments:

Drop off or send completed forms to STIC Biotech-Chem Library, Remsen Bldg.

GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: October 18, 2005, 08:22:26 ; Search time 1304 Seconds
(without alignments)
11287.047 Million cell updates/sec

Title: US-09-877-935-1
Perfect score: 8995
Sequence: 1 gatctggtgcaccaaggaca.....ctctaggctcgtccaccatg 8995

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Result No.	Score	% Query Match	Length	DB	ID	Description
	1	148.8	1.7	7208	3	US-09-166-186-107	Sequence 107, App
	2	148.8	1.7	7208	3	US-09-313-932-107	Sequence 107, App
	3	142.6	1.6	14707	4	US-09-312-762A-3	Sequence 3, Appli
c	4	140.4	1.6	3481	3	US-08-965-729A-1	Sequence 1, Appli
	5	138.2	1.5	17056	3	US-09-245-041-3	Sequence 3, Appli
	6	138.2	1.5	17056	4	US-09-358-055B-3	Sequence 3, Appli
	7	138.2	1.5	17056	4	US-09-893-238-3	Sequence 3, Appli
c	8	138	1.5	5973	3	US-09-245-041-4	Sequence 4, Appli
c	9	138	1.5	5973	4	US-09-358-055B-4	Sequence 4, Appli
c	10	138	1.5	5973	4	US-09-893-238-4	Sequence 4, Appli
c	11	136.8	1.5	2509	4	US-09-319-284-1	Sequence 1, Appli
	12	135.4	1.5	90050	3	US-09-245-041-5	Sequence 5, Appli
	13	135.4	1.5	90050	4	US-09-358-055B-5	Sequence 5, Appli
	14	135.4	1.5	90050	4	US-09-893-238-5	Sequence 5, Appli
c	15	135.2	1.5	4698	1	US-07-807-043B-5	Sequence 5, Appli
c	16	135.2	1.5	4698	1	US-08-299-849B-5	Sequence 5, Appli
c	17	135.2	1.5	4698	2	US-08-142-368A-5	Sequence 5, Appli
c	18	135.2	1.5	4698	3	US-08-967-727-5	Sequence 5, Appli
c	19	135.2	1.5	4698	3	US-08-037-230D-5	Sequence 5, Appli
c	20	135.2	1.5	4698	4	US-09-583-850-5	Sequence 5, Appli
c	21	135.2	1.5	4698	4	US-09-579-197-5	Sequence 5, Appli
c	22	135.2	1.5	4698	4	US-09-404-026-5	Sequence 5, Appli
c	23	135.2	1.5	4698	4	US-09-312-464-5	Sequence 5, Appli
	24	134	1.5	6727	3	US-08-629-643A-5	Sequence 5, Appli
	25	134	1.5	6727	3	US-09-280-799-1	Sequence 1, Appli
	26	134	1.5	6727	3	US-09-155-884-5	Sequence 5, Appli
c	27	133.4	1.5	51259	3	US-08-781-891-209	Sequence 209, App
c	28	133.4	1.5	51259	4	US-09-618-166-209	Sequence 209, App
	29	132.8	1.5	2574	4	US-09-668-313A-10	Sequence 10, Appl
c	30	132.8	1.5	17056	3	US-09-245-041-3	Sequence 3, Appli
c	31	132.8	1.5	17056	4	US-09-358-055B-3	Sequence 3, Appli
c	32	132.8	1.5	17056	4	US-09-893-238-3	Sequence 3, Appli
c	33	132.2	1.5	48974	3	US-08-920-422-17	Sequence 17, Appl
c	34	131.8	1.5	4550	3	US-09-338-907-182	Sequence 182, App
c	35	131.8	1.5	4550	3	US-09-218-207-182	Sequence 182, App
c	36	131.4	1.5	3892	2	US-08-555-723B-3	Sequence 3, Appli
c	37	131.4	1.5	3892	3	US-09-123-465-3	Sequence 3, Appli
c	38	130.8	1.5	90050	3	US-09-245-041-5	Sequence 5, Appli
c	39	130.8	1.5	90050	4	US-09-358-055B-5	Sequence 5, Appli
c	40	130.8	1.5	90050	4	US-09-893-238-5	Sequence 5, Appli
c	41	130.6	1.5	37950	3	US-09-338-907-183	Sequence 183, App
c	42	130.6	1.5	37950	3	US-09-218-207-183	Sequence 183, App
c	43	129.6	1.4	35828	3	US-09-449-218D-17	Sequence 17, Appl
c	44	129.6	1.4	35828	4	US-09-668-529A-17	Sequence 17, Appl
c	45	129.6	1.4	35828	4	US-09-668-037A-17	Sequence 17, Appl

ALIGNMENTS

RESULT 1
US-09-166-186-107
; Sequence 107, Application US/09166186A
; Patent No. 6080580
; GENERAL INFORMATION:
; APPLICANT: Baker, Brenda
; APPLICANT: Bennett, C. Frank
; APPLICANT: Butler, Madeline M.
; APPLICANT: Shanahan, William R.
; TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE MODULATION OF TNF-a EXPRESSION
; FILE REFERENCE: ISPH-0322
; CURRENT APPLICATION NUMBER: US/09/166,186A
; CURRENT FILING DATE: 1998-10-05
; NUMBER OF SEQ ID NOS: 250
; SEQ ID NO 107
; LENGTH: 7208
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (4527..4712,5225..5279,5457..5504,5799..6217)
; FEATURE:
; NAME/KEY: exon
; LOCATION: (4371)..(4712)
; FEATURE:
; NAME/KEY: intron
; LOCATION: (4713)..(5224)
; FEATURE:
; NAME/KEY: exon
; LOCATION: (5225)..(5279)
; FEATURE:
; NAME/KEY: intron
; LOCATION: (5280)..(5456)
; FEATURE:
; NAME/KEY: exon
; LOCATION: (5457)..(5504)
; FEATURE:
; NAME/KEY: intron
; LOCATION: (5505)..(5798)
; FEATURE:
; NAME/KEY: exon
; LOCATION: (5799)..(>6972)
; PUBLICATION INFORMATION:
; AUTHORS: Semon, D.
; AUTHORS: Kawashima, E.
; AUTHORS: Jongeneel, C.V.
; AUTHORS: Shakhov, A.N.
; AUTHORS: Nedospasov, S.A.
; TITLE: Nucleotide sequence of the murine TNF locus, including the
; TITLE: TNF-alpha (tumor necrosis factor) and TNF-beta (lymphotoxin)genes

```
;   JOURNAL: Nucleic Acids Res.
;   VOLUME: 15
;   ISSUE: 21
;   PAGES: 9083-9084
;   DATE: 1987-11-11
;   DATABASE ACCESSION NUMBER: Y00467 Genbank
;   DATABASE ENTRY DATE: 1993-05-11
US-09-166-186-107

  Query Match             1.7%;  Score 148.8;  DB 3;  Length 7208;
  Best Local Similarity   88.0%;  Pred. No. 4.1e-29;
  Matches  162;  Conservative    0;  Mismatches   22;  Indels    0;  Gaps    0;

Qy       4574 AGAAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGC 4633
             | ||||||| |||||   ||||||||||||||| |||||||||||||||||||||||||
Db        429 AAAAAAAAAAAAAAGCTGGGCAGTGGTGGCACACACCTTTAATCCCAGCACTTGGGAGGC 488

Qy       4634 AGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACA 4693
             ||| |||||| |||||||||||||| |||||||||||||||| |||||||||| |||||||
Db        489 AGAGGCAGGCGGATTTCTGAGTTCTAGGCCAGCCTGGTCTACAGAGTGAGTTTCAGGACA 548

Qy       4694 GCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAAC 4753
             ||||||||||||||||||||||||| | ||||||    ||||| |||| |||| ||||
Db        549 GCCAGGGCTACACAGAGAAACCCTGTCTCGAAAAAAGCAAAAAAAAAAAAAAAAAAAAA 608

Qy       4754 AAAA 4757
             ||||
Db        609 AAAA 612


RESULT 2
US-09-313-932-107
; Sequence 107, Application US/09313932A
; Patent No. 6228642
; GENERAL INFORMATION:
;  APPLICANT: Baker, Brenda
;  APPLICANT: Bennett, C. Frank
;  APPLICANT: Butler, Madeline M.
;  APPLICANT: Shanahan, William R.
;  TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE MODULATION OF TNF-
;  TITLE OF INVENTION: EXPRESSION
;  FILE REFERENCE: ISPH-0356
;  CURRENT APPLICATION NUMBER: US/09/313,932A
;  CURRENT FILING DATE: 1999-05-18
;  NUMBER OF SEQ ID NOS: 501
; SEQ ID NO 107
;   LENGTH: 7208
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (4527..4712,5225..5279,5457..5504,5799..6217)
;   FEATURE:
;   NAME/KEY: exon
;   LOCATION: (4371)..(4712)
;   FEATURE:
;   NAME/KEY: intron
;   LOCATION: (4713)..(5224)
;   FEATURE:
;   NAME/KEY: exon
;   LOCATION: (5225)..(5279)
;   FEATURE:
;   NAME/KEY: intron
;   LOCATION: (5280)..(5456)
;   FEATURE:
;   NAME/KEY: exon
;   LOCATION: (5457)..(5504)
;   FEATURE:
;   NAME/KEY: intron
;   LOCATION: (5505)..(5798)
;   FEATURE:
;   NAME/KEY: exon
;   LOCATION: (5799)..(>6972)
;  PUBLICATION INFORMATION:
;   AUTHORS: Semon, D.
;   AUTHORS: Kawashima, E.
;   AUTHORS: Jongeneel, C.V.
;   AUTHORS: Shakhov, A.N.
;   AUTHORS: Nedospasov, S.A.
;   TITLE: Nucleotide sequence of the murine TNF locus, including the
;   TITLE: TNF-alpha (tumor necrosis factor) and TNF-beta (lymphotoxin)
;   TITLE: genes
;   JOURNAL: Nucleic Acids Res.
;   VOLUME: 15
;   ISSUE: 21
;   PAGES: 9083-9084
;   DATE: 1987-11-11
;   DATABASE ACCESSION NUMBER: Y00467 Genbank
;   DATABASE ENTRY DATE: 1993-05-11
US-09-313-932-107

  Query Match             1.7%;  Score 148.8;  DB 3;  Length 7208;
  Best Local Similarity   88.0%;  Pred. No. 4.1e-29;
  Matches  162;  Conservative    0;  Mismatches   22;  Indels    0;  Gaps    0;

Qy       4574 AGAAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGC 4633
             | ||||||| |||||   ||||||||||||||| |||||||||||||||||||||||||
Db        429 AAAAAAAAAAAAAAGCTGGGCAGTGGTGGCACACACCTTTAATCCCAGCACTTGGGAGGC 488

Qy       4634 AGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACA 4693
             ||| |||||| |||||||||||||| |||||||||||||||| |||||||||| |||||||
Db        489 AGAGGCAGGCGGATTTCTGAGTTCTAGGCCAGCCTGGTCTACAGAGTGAGTTTCAGGACA 548

Qy       4694 GCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAAC 4753
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Db        549 GCCAGGGCTACACAGAGAAACCCTGTCTCGAAAAAAGCAAAAAAAAAAAAAAAAAAAAA 608

Qy       4754 AAAA 4757
             ||||
Db        609 AAAA 612


RESULT 3
US-09-312-762A-3
; Sequence 3, Application US/09312762A
; Patent No. 6552177
;  GENERAL INFORMATION:
;    APPLICANT:  MIA HOROWITZ ET AL.
;    TITLE OF INVENTION:  EH DOMAIN CONTAINING GENES AND PROTEINS
;    NUMBER OF SEQUENCES:  27
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Mark M. Friedman c/o Anthony Castorina
;      STREET:  2001 Jefferson Davis Highway, Suite 207
;      CITY:  Arlington
;      STATE:  Virginia
;      COUNTRY:  United States of America
;      ZIP:  22202
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  1.44 megabyte, 3.5" microdisk
;      COMPUTER:  Twinhead* Slimnote-890TX
;      OPERATING SYSTEM:  MS DOS version 6.2,
;      OPERATING SYSTEM:  Windows version 3.11
;      SOFTWARE:  Word for Windows version 2.0 converted to
;      SOFTWARE:  an ASCI file
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/312,762A
;      FILING DATE:
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  09/026,898
;      FILING DATE:  20 FEB 1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Friedmam, Mark M.
;      REGISTRATION NUMBER:  33,883
;      REFERENCE/DOCKET NUMBER:  916/10
;    TELECOMMUNICATION INFORMATION:
```

```
;      TELEPHONE:  972-3-5625553
;      TELEFAX:  972-3-5625554
;      TELEX:
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  14707
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
US-09-312-762A-3

  Query Match             1.6%;  Score 142.6;  DB 4;  Length 14707;
  Best Local Similarity   75.1%;  Pred. No. 3.2e-27;
  Matches  178;  Conservative    0;  Mismatches   59;  Indels    0;  Gaps    0;

Qy       4576 AAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAG 4635
              || ||  | || ||   || ||| ||||||||||||||||||||||||||||||||||||
Db      12047 AAGAAGTTTAAGGCGGGGCGGTGATGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAG 12106

Qy       4636 AAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGC 4695
              | || ||| |||||||||||||| |||||||||||||||| | ||||||||||||||||||
Db      12107 AGGCGGGCGGATTTCTGAGTTCGAGGCCAGCCTGGTCTACAAAGTGAGTTCCAGGACAGC 12166

Qy       4696 CAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAA 4755
              |||||||||||||||||||||||| |||||||||| |||||  ||| |||  |||| |
Db      12167 CAGGGCTACACAGAGAAACCCTGTCTTGAAAAACCAAAAAAAAGAAAGAAAGAAAAAAAG 12226

Qy       4756 AACAAAACCCAAACCCAAACCCAAACCTCTCATCTCTCATCTCTCTAGGCTGTGTCT 4812
              |   |||    |  ||    | |||  |  | |   |||  |  || || |
Db      12227 GTTTGAGAGCAAGTGAAGGCCTCCCCTTCTGCTGCCCCTGGACTCAGGTTTGGGTTT 12283


RESULT 4
US-08-965-729A-1/c
; Sequence 1, Application US/08965729A
; Patent No. 6200751
;  GENERAL INFORMATION:
;    APPLICANT:  Jian-Ming Gu and Charles T. Esmon
;    TITLE OF INVENTION:  ENDOTHELIUM SPECIFIC EXPRESSION
;    TITLE OF INVENTION:  REGULATED BY EPCR CONTROL ELEMENTS
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Patrea L. Pabst
;      STREET:  2800 One Atlantic Center, 1201 West Peachtree Street
;      CITY:  Atlanta
;      STATE:  GA
;      COUNTRY:  USA
;      ZIP:  30309-4530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/965,729A
;      FILING DATE:  07-NOV-1997
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/030,718
;      FILING DATE:  08-NOV-1997
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  OMRF 164 PCT
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-873-8794
;      TELEFAX:  404-873-8795
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3481 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  misc_feature
;      OTHER INFORMATION:  /note= "Nucleotides 2270 through 2840 are a
;      OTHER INFORMATION:  large endothelial specific element"; murine
;    FEATURE:
;      NAME/KEY:  misc_feature
;      OTHER INFORMATION:  /note= "Nucleotides 2990 through 3061 are a
;      OTHER INFORMATION:  serum response element"; murine
;    FEATURE:
;      NAME/KEY:  misc_feature
;      OTHER INFORMATION:  /note= "Nucleotides 3007 through 3014 are a
;      OTHER INFORMATION:  thrombin responsive element"; murine
;    FEATURE:
;      NAME/KEY:  misc_feature
;      OTHER INFORMATION:  /note= "Nucleotides 3130 through 3350 are an
;      OTHER INFORMATION:  endothelial specific element"; murine
US-08-965-729A-1

  Query Match             1.6%;  Score 140.4;  DB 3;  Length 3481;
  Best Local Similarity   74.8%;  Pred. No. 4.8e-27;
  Matches  202;  Conservative    0;  Mismatches   66;  Indels    2;  Gaps    2;

Qy       4564 TTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCA 4623
              ||   |||||| |||||||  |||  || |  ||||||||||||||||||||||||||||
Db       2563 TTCAACTTTAAAAAAAAAAAAAAAAGCCGGGCGTGGTGGCACACGCCTTTAATCCCAGCA 2504

Qy       4624 CTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAG 4683
              |||||||||||| |||||  |||||||||||  |||||||||| ||| ||||||||
Db       2503 CTTGGGAGGCAGAGGCAGGTGGATTTCTGAGTTTGAGGCCAGCCTGGCCTACAGAGTGAG 2444

Qy       4684 TTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAA-AAACAAAA 4742
              ||||||||||||||||||||||||| |||||||||  |  |||||| | || ||| | ||
Db       2443 -TCCAGGACAGCCAGGGCTACACAGAAAAACCCTGTCTCAAAAAACAAAAACAAAAACAA 2385

Qy       4743 CAAAACAAAACAAAACAAAACCCAAACCCAAACCCAAACCTCTCATCTCTCATCTCTCTA 4802
              ||||||||||||||||      ||| |||||   |||||    |  |  |     | |
Db       2384 AAAAACAAAACAAAACACCCAAAAAAACCAAAAAAAAAACCCAAAAAACCATAAAAACCCA 2325

Qy       4803 GGCTGTGTCTGTCTAGGTGGTAGAGTTTGG 4832
              | | ||||  |||       ||| ||||
Db       2324 AACAGGGTCTTACTATCCATATGAGGTTGG 2295


RESULT 5
US-09-245-041-3
; Sequence 3, Application US/09245041
; Patent No. 6274339
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT: Nagle, D.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT
;  TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY
;  FILE REFERENCE: 7853-136
;  CURRENT APPLICATION NUMBER: US/09/245,041
;  CURRENT FILING DATE: 1999-02-05
;  EARLIER APPLICATION NUMBER: 60/093,630
;  EARLIER FILING DATE: 1998-07-21
;  EARLIER APPLICATION NUMBER: 60/104,978
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 131
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 3
;   LENGTH: 17056
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-245-041-3

  Query Match             1.5%;  Score 138.2;  DB 3;  Length 17056;
  Best Local Similarity   79.2%;  Pred. No. 5.7e-26;
  Matches  164;  Conservative    0;  Mismatches   43;  Indels    0;  Gaps    0;
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Qy       4550 CTATAGAGCGAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGC 4609
Db      10295 CCAAACACCCATACAAACCAATTTTTAAAAAAATATTCGAGCCGGCGTGGTGGCGCACGC 10354

Qy       4610 CTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTG 4669
Db      10355 CTTTAATCCCAGCACTCGGGAGACAGAGGCAGGTGGATTTCTGAGTTCGAGGCCAGCGTG 10414

Qy       4670 GTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAAC 4729
Db      10415 GTCTACAGAGTGAGTTCCAGGACAGCCAGGGCTGCACAGAGAAACCCTGTCTCGAAAAAC 10474

Qy       4730 CAGAAAAACAAAACAAAACAAAACAAA 4756
Db      10475 CAAAAAAAAAAAAAAAAAAAAAAATA 10501


RESULT 6
US-09-358-055B-3
; Sequence 3, Application US/09358055B
; Patent No. 6713277
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT:  Nagle, D.L.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND
;  TITLE OF INVENTION:  TREATMENT OF BODY WEIGHT DISORDERS INCLUDING
;  TITLE OF INVENTION:  OBESITY
;  FILE REFERENCE: 7853-151
;  CURRENT APPLICATION NUMBER: US/09/358,055B
;  CURRENT FILING DATE:  1999-07-21
;  PRIOR APPLICATION NUMBER: 09/245,041
;  PRIOR FILING DATE: 1999-02-05
;  NUMBER OF SEQ ID NOS: 153
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 3
;   LENGTH: 17056
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-358-055B-3

  Query Match             1.5%;  Score 138.2;  DB 4;  Length 17056;
  Best Local Similarity   79.2%;  Pred. No. 5.7e-26;
  Matches  164;  Conservative    0;  Mismatches   43;  Indels    0;  Gaps    0;

Qy       4550 CTATAGAGCGAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGC 4609
Db      10295 CCAAACACCCATACAAACCAATTTTTAAAAAAATATTCGAGCCGGCGTGGTGGCGCACGC 10354

Qy       4610 CTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTG 4669
Db      10355 CTTTAATCCCAGCACTCGGGAGACAGAGGCAGGTGGATTTCTGAGTTCGAGGCCAGCGTG 10414

Qy       4670 GTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAAC 4729
Db      10415 GTCTACAGAGTGAGTTCCAGGACAGCCAGGGCTGCACAGAGAAACCCTGTCTCGAAAAAC 10474

Qy       4730 CAGAAAAACAAAACAAAACAAAACAAA 4756
Db      10475 CAAAAAAAAAAAAAAAAAAAAAAATA 10501


RESULT 7
US-09-893-238-3
; Sequence 3, Application US/09893238
; Patent No. 6727348
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT:  Nagle, D.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND
;  TITLE OF INVENTION:  DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY
;  FILE REFERENCE: 7853-237
;  CURRENT APPLICATION NUMBER: US/09/893,238
;  CURRENT FILING DATE:  2001-06-27
;  PRIOR APPLICATION NUMBER: 09/245,041
;  PRIOR FILING DATE: 1999-02-05
;  PRIOR APPLICATION NUMBER: 60/093,630
;  PRIOR FILING DATE: 1998-07-21
;  PRIOR APPLICATION NUMBER: 60/104,978
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 129
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 3
;   LENGTH: 17056
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-893-238-3

  Query Match             1.5%;  Score 138.2;  DB 4;  Length 17056;
  Best Local Similarity   79.2%;  Pred. No. 5.7e-26;
  Matches  164;  Conservative    0;  Mismatches   43;  Indels    0;  Gaps    0;

Qy       4550 CTATAGAGCGAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGC 4609
Db      10295 CCAAACACCCATACAAACCAATTTTTAAAAAAATATTCGAGCCGGCGTGGTGGCGCACGC 10354

Qy       4610 CTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTG 4669
Db      10355 CTTTAATCCCAGCACTCGGGAGACAGAGGCAGGTGGATTTCTGAGTTCGAGGCCAGCGTG 10414

Qy       4670 GTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAAC 4729
Db      10415 GTCTACAGAGTGAGTTCCAGGACAGCCAGGGCTGCACAGAGAAACCCTGTCTCGAAAAAC 10474

Qy       4730 CAGAAAAACAAAACAAAACAAAACAAA 4756
Db      10475 CAAAAAAAAAAAAAAAAAAAAAAATA 10501


RESULT 8
US-09-245-041-4/c
; Sequence 4, Application US/09245041
; Patent No. 6274339
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT: Nagle, D.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT
;  TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY
;  FILE REFERENCE: 7853-136
;  CURRENT APPLICATION NUMBER: US/09/245,041
;  CURRENT FILING DATE: 1999-02-05
;  EARLIER APPLICATION NUMBER: 60/093,630
;  EARLIER FILING DATE: 1998-07-21
;  EARLIER APPLICATION NUMBER: 60/104,978
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 131
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 4
;   LENGTH: 5973
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-245-041-4

  Query Match             1.5%;  Score 138;  DB 3;  Length 5973;
  Best Local Similarity   83.9%;  Pred. No. 3.1e-26;
  Matches  156;  Conservative    0;  Mismatches   30;  Indels    0;  Gaps    0;

Qy       4572 TAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAG 4631
Db       5564 TAAGAAAATCTTTTAGGCTGGGTGGTGGTGGCACACACCTTTAATCCCAGCATTTGGGAG 5505

Qy       4632 GCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGA 4691
Db       5504 GCAGAGGCAGGCAGATTTCTGAGTTCGAGGCCAGCCTGGTCTACAGAGTGAGTTCCAGGA 5445

Qy       4692 CAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAA 4751
```

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              |||||||||||| |||||||||||||||| |||||||||| |||   ||||  ||  | |
Db       5444 CAGCCAGGGCTATACAGAGAAACCCTGTCTTGAAAAACCGAAAATGAAAAAAGAAGAAGA 5385

Qy       4752 ACAAAA 4757
              | || |
Db       5384 AGAAGA 5379


RESULT 9
US-09-358-055B-4/c
; Sequence 4, Application US/09358055B
; Patent No. 6713277
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT:  Nagle, D.L.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND
;  TITLE OF INVENTION:  TREATMENT OF BODY WEIGHT DISORDERS INCLUDING
;  TITLE OF INVENTION:  OBESITY
;  FILE REFERENCE: 7853-151
;  CURRENT APPLICATION NUMBER: US/09/358,055B
;  CURRENT FILING DATE:  1999-07-21
;  PRIOR APPLICATION NUMBER: 09/245,041
;  PRIOR FILING DATE: 1999-02-05
;  NUMBER OF SEQ ID NOS: 153
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 4
;   LENGTH: 5973
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-358-055B-4

  Query Match             1.5%;  Score 138;  DB 4;  Length 5973;
  Best Local Similarity   83.9%;  Pred. No. 3.1e-26;
  Matches  156;  Conservative    0;  Mismatches   30;  Indels    0;  Gaps    0;

Qy       4572 TAAGAAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAG 4631
              ||||||||    |  | ||   |  |||||||||||| |||||||||||||| |||||||
Db       5564 TAAGAAAATCTTTTAGGCTGGGTGGTGGTGGCACACACCTTTAATCCCAGCATTTGGGAG 5505

Qy       4632 GCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGA 4691
              ||||| |||||||||||||||||| |||||||||||||| |||||||||||||||
Db       5504 GCAGAGGCAGGCAGATTTCTGAGTTCGAGGCCAGCCTGGTCTACAGAGTGAGTTCCAGGA 5445

Qy       4692 CAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAA 4751
              |||||||||||| |||||||||||||||| |||||||||  |||   ||||  ||  | |
Db       5444 CAGCCAGGGCTATACAGAGAAACCCTGTCTTGAAAAACCGAAAATGAAAAAAGAAGAAGA 5385

Qy       4752 ACAAAA 4757
              | || |
Db       5384 AGAAGA 5379


RESULT 10
US-09-893-238-4/c
; Sequence 4, Application US/09893238
; Patent No. 6727348
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT:  Nagle, D.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND
;  TITLE OF INVENTION:  DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY
;  FILE REFERENCE: 7853-237
;  CURRENT APPLICATION NUMBER: US/09/893,238
;  CURRENT FILING DATE:  2001-06-27
;  PRIOR APPLICATION NUMBER: 09/245,041
;  PRIOR FILING DATE: 1999-02-05
;  PRIOR APPLICATION NUMBER: 60/093,630
;  PRIOR FILING DATE: 1998-07-21
;  PRIOR APPLICATION NUMBER: 60/104,978
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 129
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 4
;   LENGTH: 5973
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-893-238-4

  Query Match             1.5%;  Score 138;  DB 4;  Length 5973;
  Best Local Similarity   83.9%;  Pred. No. 3.1e-26;
  Matches  156;  Conservative    0;  Mismatches   30;  Indels    0;  Gaps    0;

Qy       4572 TAAGAAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAG 4631
              ||||||||    |  | ||   |  |||||||||||| |||||||||||||| |||||||
Db       5564 TAAGAAAATCTTTTAGGCTGGGTGGTGGTGGCACACACCTTTAATCCCAGCATTTGGGAG 5505

Qy       4632 GCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGA 4691
              ||||| |||||||||||||||||| |||||||||||||| |||||||||||||||
Db       5504 GCAGAGGCAGGCAGATTTCTGAGTTCGAGGCCAGCCTGGTCTACAGAGTGAGTTCCAGGA 5445

Qy       4692 CAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAA 4751
              |||||||||||| |||||||||||||||| |||||||||  |||   ||||  ||  | |
Db       5444 CAGCCAGGGCTATACAGAGAAACCCTGTCTTGAAAAACCGAAAATGAAAAAAGAAGAAGA 5385

Qy       4752 ACAAAA 4757
              | || |
Db       5384 AGAAGA 5379


RESULT 11
US-09-319-284-1/c
; Sequence 1, Application US/09319284A
; Patent No. 6524815
; GENERAL INFORMATION:
;  APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE-CEA
;  APPLICANT: HUBER, Philippe
;  APPLICANT: LAURENT, Monique
;  APPLICANT: GORY, Sylvie
;  TITLE OF INVENTION: VE CADHERIN PROMOTER AND ITS USES
;  FILE REFERENCE: 45636-5020-US
;  CURRENT APPLICATION NUMBER: US/09/319,284A
;  CURRENT FILING DATE: 1999-08-27
;  EARLIER APPLICATION NUMBER: PCT/FR97/02178
;  EARLIER FILING DATE: 1997-12-02
;  EARLIER APPLICATION NUMBER: FR 9614801
;  EARLIER FILING DATE: 1996-12-03
;  NUMBER OF SEQ ID NOS: 3
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
;   LENGTH: 2509
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-319-284-1

  Query Match             1.5%;  Score 136.8;  DB 4;  Length 2509;
  Best Local Similarity   87.2%;  Pred. No. 3.6e-26;
  Matches  150;  Conservative    0;  Mismatches   22;  Indels    0;  Gaps    0;

Qy       4590 CCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTT 4649
              ||||  |||||||| |||||||||||||||||||| |||||||||| |||||  |||||
Db       1657 CCAGGCGTGGTGGCGCACGCCTTTAATCCCAGCACTCGGGAGGCAGAGGCAGGTGGATTT 1598

Qy       4650 CTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGA 4709
              |||||||| |||||||||||||||| |||||| |||||||| |||||||||| |||||
Db       1597 CTGAGTTCGAGGCCAGCCTGGTCTACAGAGTGTGTTCCAGGATAGCCAGGGCTATACAGA 1538

Qy       4710 GAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAA 4761
              |||||||||| | |||||||||| ||||| | || |||| |||| || | |||
Db       1537 GAAACCCTGTCTCGAAAAACCAAAAAAAAAGAAAAAAAAAAAAAAAGAAAAA 1486


RESULT 12
US-09-245-041-5
; Sequence 5, Application US/09245041
; Patent No. 6274339
; GENERAL INFORMATION:
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;  APPLICANT: Moore, K.
;  APPLICANT: Nagle, D.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT
;  TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY
;  FILE REFERENCE: 7853-136
;  CURRENT APPLICATION NUMBER: US/09/245,041
;  CURRENT FILING DATE: 1999-02-05
;  EARLIER APPLICATION NUMBER: 60/093,630
;  EARLIER FILING DATE: 1998-07-21
;  EARLIER APPLICATION NUMBER: 60/104,978
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 131
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 5
;   LENGTH: 90050
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-245-041-5

  Query Match             1.5%;  Score 135.4;  DB 3;  Length 90050;
  Best Local Similarity   87.6%;  Pred. No. 1e-24;
  Matches  148;  Conservative    0;  Mismatches   21;  Indels    0;  Gaps    0;

Qy       4593 GCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTG 4652
             | |||||||||||||||||||||||||||||||||||||||||||||||||| |||||  ||||||||
Db      16335 GTAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAGGCAGGTGGATTTCTG 16394

Qy       4653 AGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAA 4712
             ||||| || |||||||||| || ||||||||||||||||||||||||| ||||||||||||
Db      16395 AGTTCGAGACCAGCCTGGTTTACAGAGTGAGTTCCAGGACAGCCAGGGATACACAGAGAA 16454

Qy       4713 ACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAA 4761
              ||||||| | ||||| | |  |||  |||||||||||||  || | |||
Db      16455 ACCCTGTCTCGAAAAGCAAACAAACAAAAACAAAACAAACAAACAAAAA 16503


RESULT 13
US-09-358-055B-5
; Sequence 5, Application US/09358055B
; Patent No. 6713277
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT:  Nagle, D.L.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND
;  TITLE OF INVENTION:  TREATMENT OF BODY WEIGHT DISORDERS INCLUDING
;  TITLE OF INVENTION:  OBESITY
;  FILE REFERENCE: 7853-151
;  CURRENT APPLICATION NUMBER: US/09/358,055B
;  CURRENT FILING DATE:  1999-07-21
;  PRIOR APPLICATION NUMBER: 09/245,041
;  PRIOR FILING DATE: 1999-02-05
;  NUMBER OF SEQ ID NOS: 153
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 5
;   LENGTH: 90050
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-358-055B-5

  Query Match             1.5%;  Score 135.4;  DB 4;  Length 90050;
  Best Local Similarity   87.6%;  Pred. No. 1e-24;
  Matches  148;  Conservative    0;  Mismatches   21;  Indels    0;  Gaps    0;

Qy       4593 GCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTG 4652
             | |||||||||||||||||||||||||||||||||||||||||||||||||| |||||  ||||||||
Db      16335 GTAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAGGCAGGTGGATTTCTG 16394

Qy       4653 AGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAA 4712
             ||||| || |||||||||| || ||||||||||||||||||||||||| ||||||||||||
Db      16395 AGTTCGAGACCAGCCTGGTTTACAGAGTGAGTTCCAGGACAGCCAGGGATACACAGAGAA 16454

Qy       4713 ACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAA 4761
              ||||||| | ||||| | |  |||  |||||||||||||  || | |||
Db      16455 ACCCTGTCTCGAAAAGCAAACAAACAAAAACAAAACAAACAAACAAAAA 16503


RESULT 14
US-09-893-238-5
; Sequence 5, Application US/09893238
; Patent No. 6727348
; GENERAL INFORMATION:
;  APPLICANT: Moore, K.
;  APPLICANT:  Nagle, D.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND
;  TITLE OF INVENTION:  DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY
;  FILE REFERENCE: 7853-237
;  CURRENT APPLICATION NUMBER: US/09/893,238
;  CURRENT FILING DATE:  2001-06-27
;  PRIOR APPLICATION NUMBER: 09/245,041
;  PRIOR FILING DATE: 1999-02-05
;  PRIOR APPLICATION NUMBER: 60/093,630
;  PRIOR FILING DATE: 1998-07-21
;  PRIOR APPLICATION NUMBER: 60/104,978
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 129
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 5
;   LENGTH: 90050
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-893-238-5

  Query Match             1.5%;  Score 135.4;  DB 4;  Length 90050;
  Best Local Similarity   87.6%;  Pred. No. 1e-24;
  Matches  148;  Conservative    0;  Mismatches   21;  Indels    0;  Gaps    0;

Qy       4593 GCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTG 4652
             | |||||||||||||||||||||||||||||||||||||||||||||||||| |||||  ||||||||
Db      16335 GTAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAGGCAGGTGGATTTCTG 16394

Qy       4653 AGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAA 4712
             ||||| || |||||||||| || ||||||||||||||||||||||||| ||||||||||||
Db      16395 AGTTCGAGACCAGCCTGGTTTACAGAGTGAGTTCCAGGACAGCCAGGGATACACAGAGAA 16454

Qy       4713 ACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAA 4761
              ||||||| | ||||| | |  |||  |||||||||||||  || | |||
Db      16455 ACCCTGTCTCGAAAAGCAAACAAACAAAAACAAAACAAACAAACAAAAA 16503


RESULT 15
US-07-807-043B-5/c
; Sequence 5, Application US/07807043B
; Patent No. 5342774
;  GENERAL INFORMATION:
;    APPLICANT:  Boon, Thierry, Van den Eynde, Beno t
;    TITLE OF INVENTION:  Tumor Rejection Antigen Precursors, Tumor
;    TITLE OF INVENTION:  Rejection Antigens and Uses Thereof
;    NUMBER OF SEQUENCES:  16
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Felfe & Lynch
;      STREET:  805 Third Avenue
;      CITY:  New York City
;      STATE:  New York
;      ZIP:  10022
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette, 5.25 inch, 360 kb storage
;      COMPUTER:  IBM
;      OPERATING SYSTEM:  PC-DOS
;      SOFTWARE:  Wordperfect
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/807,043B
;      FILING DATE:  19911212
;      CLASSIFICATION:  424
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  07/764,364
;      FILING DATE:  23-SEPTEMBER-1991
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; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/728,838
; FILING DATE: 9-JULY-1991
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/705,702
; FILING DATE: 23-May-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Hanson, No. 5342774man D.
; REGISTRATION NUMBER: 30,946
; REFERENCE/DOCKET NUMBER: LUD 253.3
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 688-9200
; TELEFAX: (212) 838-3884
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 4698 base pairs
; TYPE: NUCLEIC ACID
; STRANDEDNESS: singular
; TOPOLOGY: linear
; MOLECULE TYPE: genomic DNA
US-07-807-043B-5

Query Match 1.5%; Score 135.2; DB 1; Length 4698;
Best Local Similarity 86.6%; Pred. No. 1.5e-25;
Matches 149; Conservative 0; Mismatches 23; Indels 0; Gaps 0;

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Qy       4593 GCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTG 4652
             ||||| ||| || ||||||||||||||| |||||||||||| ||||||||||||||||
Db       1706 GCAGTCCTGGTGCAAGCCTTTAATCCCAGCATTTGGGAGGCAGAGGCAGGCAGATTTCTG 1647

Qy       4653 AGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAA 4712
             |||| |||||||||||||||| ||||||||| |||||||||||||||| |||| ||||||
Db       1646 AGTTTGAGGCCAGCCTGGTCTACAGAGTGAGTGCCAGGACAGCCAGGGATACAAAGAGAA 1587

Qy       4713 ACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAAACC 4764
             ||||||| | ||||||||| ||||| |||| |||| |||| |||| || ||
Db       1586 ACCCTGTCTCGAAAAACCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAGGCC 1535
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Job time : 1306 secs

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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: October 18, 2005, 08:22:27 ; Search time 5397 Seconds
(without alignments)
11685.034 Million cell updates/sec

Title: US-09-877-935-1
Perfect score: 8995
Sequence: 1 gatctggtgcaccaaggaca.....ctctaggctcgtccaccatg 8995

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 8766186 seqs, 3505510206 residues

Total number of hits satisfying chosen parameters: 17532372

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
6: /cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
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11: /cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
12: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
13: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
14: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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	1	8995	100.0	8995	9	US-09-877-935-1	Sequence 1, Appli
c	2	170.4	1.9	493631	13	US-10-087-192-205	Sequence 205, App
	3	166.6	1.9	63502	13	US-10-087-192-271	Sequence 271, App
c	4	165	1.8	33488	17	US-10-085-117-235	Sequence 235, App
c	5	164	1.8	31632	17	US-10-034-650-1	Sequence 1, Appli
	6	161.4	1.8	21781	19	US-10-322-696-25	Sequence 25, Appl
	7	161.4	1.8	54648	17	US-10-085-117-289	Sequence 289, App
c	8	161	1.8	111836	19	US-10-322-281-51	Sequence 51, Appl
	9	158.8	1.8	27383	19	US-10-322-696-133	Sequence 133, App
	10	158	1.8	26345	13	US-10-087-192-1705	Sequence 1705, Ap
c	11	157.6	1.8	193853	13	US-10-087-192-1663	Sequence 1663, Ap
c	12	156.2	1.7	93329	19	US-10-322-281-817	Sequence 817, App
	13	156	1.7	24495	19	US-10-367-094-142	Sequence 142, App
	14	155.6	1.7	31400	13	US-10-087-192-907	Sequence 907, App
	15	155	1.7	94381	11	US-09-997-722-31	Sequence 31, Appl
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c	17	154.4	1.7	29222	17	US-10-085-117-349	Sequence 349, App
c	18	153.4	1.7	96389	18	US-10-052-482-181	Sequence 181, App
c	19	153	1.7	72821	21	US-10-461-862-149	Sequence 149, App
	20	152.8	1.7	247461	19	US-10-322-281-131	Sequence 131, App
	21	152.4	1.7	197775	13	US-10-087-192-853	Sequence 853, App
	22	152.2	1.7	170279	19	US-10-388-838-1	Sequence 1, Appli
	23	152.2	1.7	215980	10	US-09-972-546-16	Sequence 16, Appl
	24	152.2	1.7	215980	21	US-10-735-256-16	Sequence 16, Appl
c	25	151.2	1.7	122859	13	US-10-087-192-37	Sequence 37, Appl
c	26	151	1.7	46677	13	US-10-087-192-943	Sequence 943, App
	27	151	1.7	116585	13	US-10-087-192-133	Sequence 133, App
	28	150.6	1.7	1445	17	US-10-461-093-17	Sequence 17, Appl
	29	150.4	1.7	53828	19	US-10-322-281-369	Sequence 369, App
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	32	150	1.7	53158	13	US-10-087-192-1963	Sequence 1963, Ap
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	34	149.8	1.7	49088	13	US-10-087-192-13	Sequence 13, Appl
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c	36	149.8	1.7	263852	20	US-10-812-232-6	Sequence 6, Appli
	37	149.6	1.7	224112	19	US-10-367-094-80	Sequence 80, Appl
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c	40	149	1.7	29956	11	US-09-997-722-229	Sequence 229, App
c	41	149	1.7	44990	18	US-10-052-482-217	Sequence 217, App
c	42	149	1.7	83888	13	US-10-087-192-541	Sequence 541, App
c	43	149	1.7	84252	19	US-10-322-281-563	Sequence 563, App
	44	149	1.7	158405	14	US-10-175-523-86	Sequence 86, Appl
	45	149	1.7	158405	24	US-11-099-266-86	Sequence 86, Appl

ALIGNMENTS

RESULT 1
US-09-877-935-1
; Sequence 1, Application US/09877935
; Patent No. US20020102705A1
; GENERAL INFORMATION:
; APPLICANT: Pinto, Daniel
; APPLICANT: Robine, Sylvie
; APPLICANT: Jaisser, Frederic
; APPLICANT: Louvard, Daniel
; TITLE OF INVENTION: REGULATORY SEQUENCES OF THE MOUSE VILLIN GENE - USE IN TRANSGENES
; FILE REFERENCE: 13294-002001
; CURRENT APPLICATION NUMBER: US/09/877,935
; CURRENT FILING DATE: 2001-06-08
; PRIOR APPLICATION NUMBER: PCT/EP 98/08009
; PRIOR FILING DATE: 1998-12-09
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 8995
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; FEATURE:
; NAME/KEY: intron
; LOCATION: (3489)..(8981)
; NAME/KEY: exon
; LOCATION: (3443)..(3487)
; OTHER INFORMATION: exon 1
US-09-877-935-1

Query Match 100.0%; Score 8995; DB 9; Length 8995;

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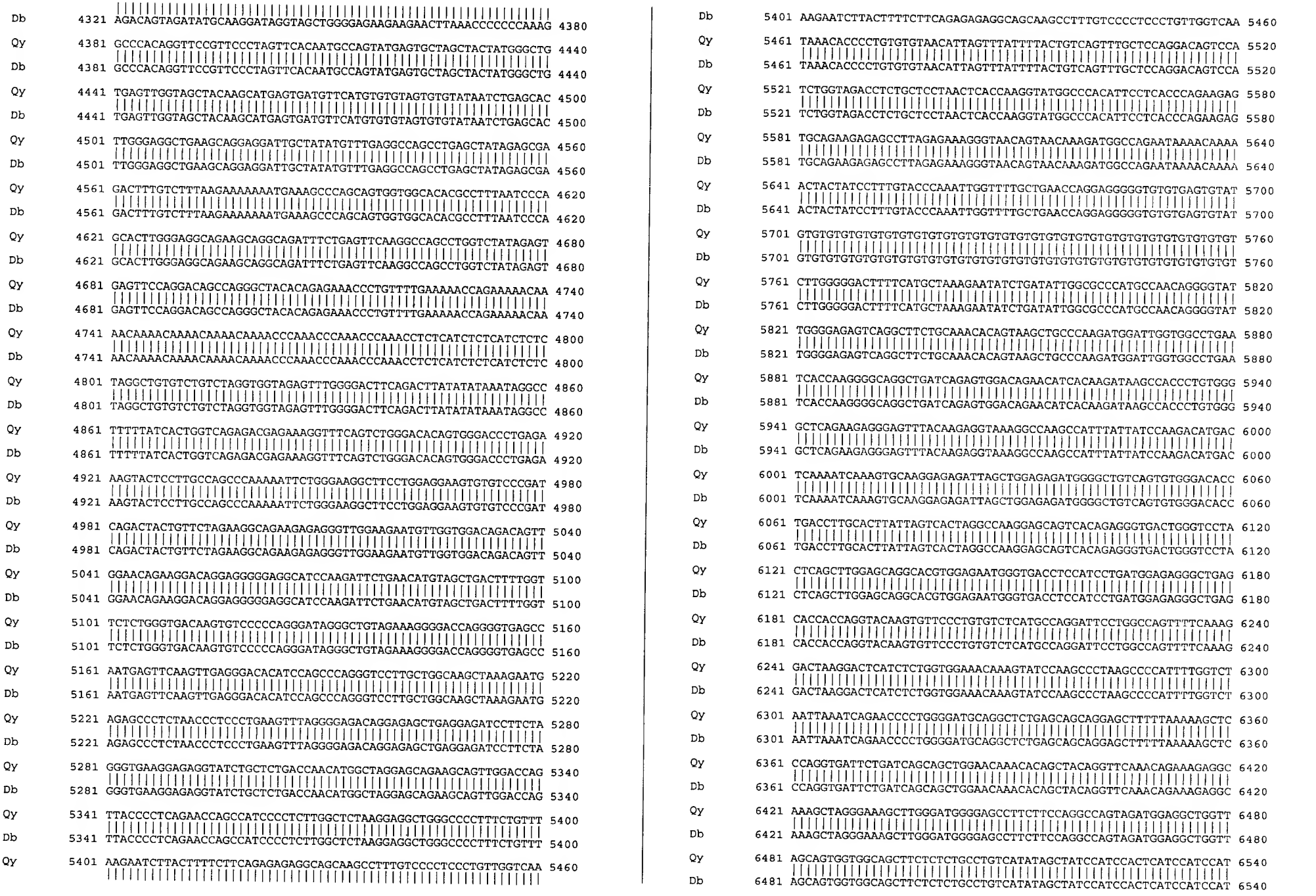
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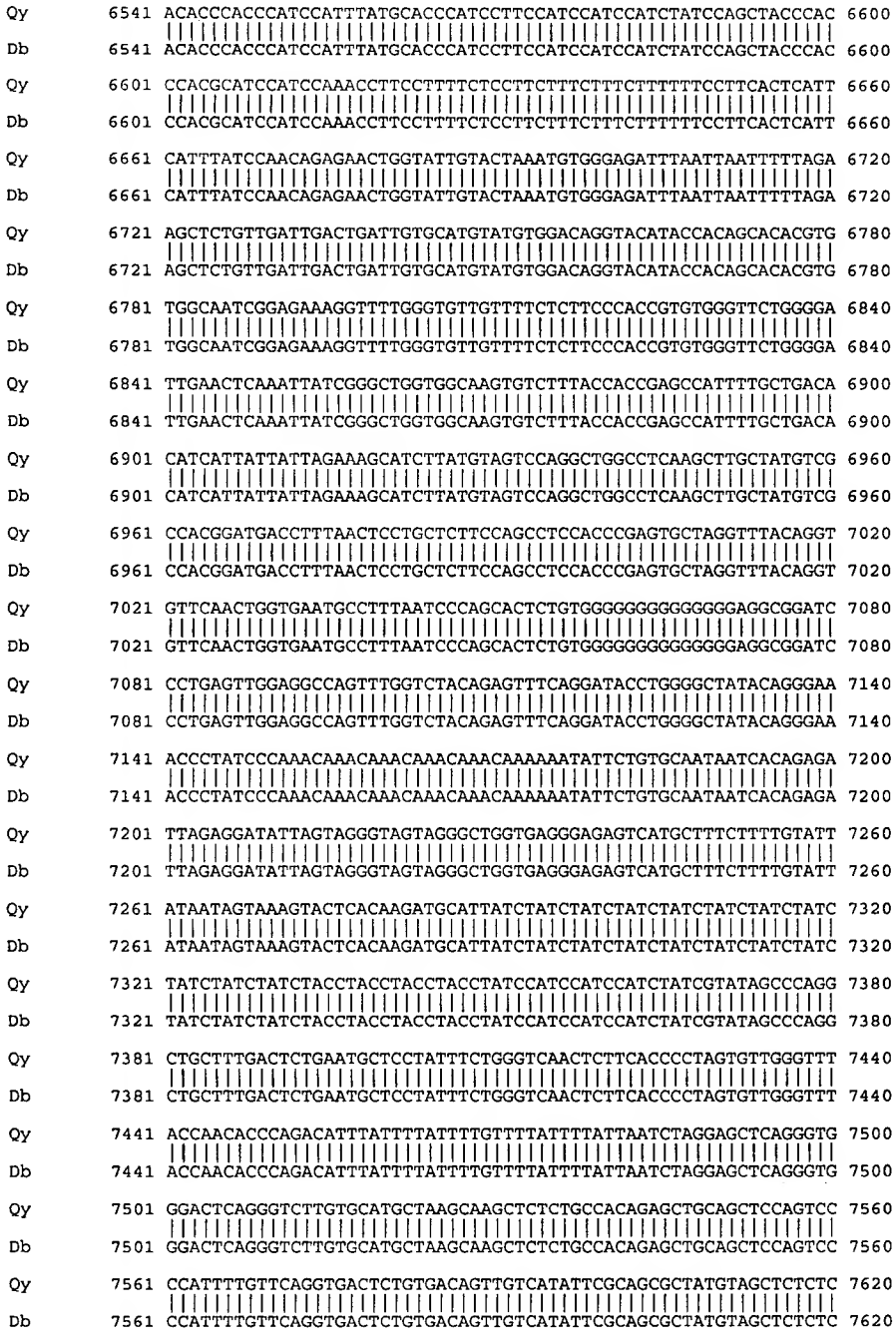
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; Publication No. US20020182586A1
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; APPLICANT: Morris, David W.
; APPLICANT: Engelhard, Eric K.
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
; TITLE OF INVENTION: CANCER
; FILE REFERENCE: 529452000122
; CURRENT APPLICATION NUMBER: US/10/087,192
; CURRENT FILING DATE: 2002-03-01
; PRIOR APPLICATION NUMBER: US 09/747,377
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
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Db      28871 ACTCCAGTTTTAAGAGATCGAATGCTCTCTTCTGAGGCCCCAGGGGCACTGCACACACAT 28812

Qy       2012 TACACA-GACATACATGCAGGCAAAACAC-CCATACACACATAAATAAATAAGGAAACTT 2069
Db      28811 GACATAGGACACACATGTAGAAAAAACACATAAAATAAGAATAACTCACTCTGTAGACCA 28752

Qy       2070 AAAAGGTGCATGTGTTGGTAAACATTGTGCTTACACATGCTGATTGAAGACATGTACAAC 2129
Db      28751 GACTGGCCTCGAACTCAGAAATCCACCTGCCTCTGCTTCCCAAGTGCTGGGATTAAAGGC 28692

Qy       2130 GCACAC----------ACTGAAGAGGGATCTGGGGCTGGAGA 2161
```

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Db      28691 GTGCGCCACCACTGCTCAGCTTAAAAAGTTCTCTTAAGAACAGTAATGACGGGCTGGAGA 28632

Qy       2162 GATGGCTCAGCGGTTAAGAGCACTGACTGCTCTTCCGAAGGAAGGTCCTGAGTTCAAATC 2221
Db      28631 GATGGCTCAGTGGTTAAGAGCACCGACTGCTCTTCC----GAAGATCCTGAGTTCAAATC 28576

Qy       2222 CTAGCAACCACATGGTGGCTCACAACCATCCATAATGAGATCTGACACCCTCTTCTGGTG 2281
Db      28575 CCAGCAACCACATGGTAGCTCACAACCATCCATAATGAGATCTGACGCCCTCTTCTGGTG 28516

Qy       2282 CATCTGAAGACAGCTGCAGAGCTACAGTGTACTTAGATATACTAATAAATAAATCTT--T 2339
Db      28515 TGTCTGAAGAC--------AGTTACAGTGTACTTAGATATAATAATAAATAAATCTTAAA 28464

Qy       2340 TTTTAAAAAAATGAAGAGGGATCTGAGACACCTCAAAAGAGATTATGAGCAGTGACTCAC 2399
Db      28463 AAAAAAAAAAAAGAACAGTAATGACGATGATGATGATAAAGAGTCTTTGCTGTTACTCAA 28404

Qy       2400 GGGTGATTATCTATCCTGGA 2419
Db      28403 CTGAAATAATTTCTTAGGGA 28384
```

RESULT 3
US-10-087-192-271
; Sequence 271, Application US/10087192
; Publication No. US20020182586A1
; GENERAL INFORMATION:
; APPLICANT: Morris, David W.
; APPLICANT: Engelhard, Eric K.
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
; TITLE OF INVENTION: CANCER
; FILE REFERENCE: 529452000122
; CURRENT APPLICATION NUMBER: US/10/087,192
; CURRENT FILING DATE: 2002-03-01
; PRIOR APPLICATION NUMBER: US 09/747,377
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 2059
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 271
; LENGTH: 63502
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(63502)
; OTHER INFORMATION: n = A,T,C or G
US-10-087-192-271

Query Match 1.9%; Score 166.6; DB 13; Length 63502;
Best Local Similarity 64.6%; Pred. No. 3.5e-34;
Matches 330; Conservative 0; Mismatches 169; Indels 12; Gaps 5;

```
Qy       1857 GGGAGAGATGGCTCAGCTTCCAGGAGCACTTGCTGCTCTTGCAGAGGACCTAGATTCAGT 1916
Db       2258 GAGAGAGAGAGAGAGCTGTCAGTGGTTAAGAGCTGCTATTGCGGAGGACCTTGGTTCGGT 2317

Qy       1917 TCCCAGGACTCATATGGTGGCTCACAGCCATCTGTAAATCCAGTTCCAGAGGGTTCCACA 1976
Db       2318 TCCTAGCACCTACATGG-GGTTCACAATCATCCCTAACTCCAGTTCCAT-GGGATCAGCT 2375

Qy       1977 CCCTCTTCTGGCCTCCACAGGCAC----CACATACATAGTACACAGACATACATGCAGGC 2032
Db       2376 AGCCCTTCTGACCTCTGCAGGCACTATGCATACATGTGGTGCACATACATACATGTAAGC 2435

Qy       2033 AAAACACCCATACACACATAAATAAATAAGGAAACTTAAAAGGTGCATGTGTTGGTAAAC 2092
Db       2436 AAAACACATGAAGAAAAATGAATACATCTTGTAAATCTTTAAAGTCATTAATT--TTAAA 2493

Qy       2093 ATTGTGCTTACACATGCTGATTGAAGACATGTACAACGCACACACTGAAGAGGGATCTGG 2152
```

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                  | | | |  |    |  | || ||     |    | | ||       ||      | |||
Db       2494 AGTATCCAAATGATTCTTTATGGATTCTTTTGGTATCTCAAGTGAAAAAATCAATTTTGG 2553

Qy       2153 GGCTGGAGAGATGGCTCAGCGGTTAAGAGCACTGACTGCTCTTCCGAAGGAAGGTCCTGA 2212
              |||||||||||||||| |  ||||||||||||||||||||||||     |||||  |||
Db       2554 GGCTGGAGAGATGGCTCCGTAGTTAAGAGCACTGACTGCTCTTCCAG----AGGTCATGA 2609

Qy       2213 GTTCAAATCCTAGCAACCACATGGTGGCTCACAACCATCCATAATGAGATCTGACACCCT 2272
              |||||| ||| |||| ||||||||||||||||||||||  |||||  |||||||  ||||
Db       2610 GTTCAATTCCCAGCAGCCACATGGTGGCTCACAACCATCTGTAATGGGATCTGATGCCCT 2669

Qy       2273 CTTCTGGTGCATCTGAAGACAGCTGCAGAGCTACAGTGTACTTAGATATACTAATAAATA 2332
              ||||||||| ||||| |||||||| ||| |      |     |     | |  ||||||||
Db       2670 CTTCTGGTGTATCTGCAGACAGCTACAGTGTATTCATCATGTATATAAAATAAATAAATA 2729

Qy       2333 AATCTTTTTTTAAAAAAATGAAGAGGGATCT 2363
              |   ||||  | |||||  || ||  || |
Db       2730 CAATCTTTTAAAGAAAAAGAAAAAGCAATTT 2760


RESULT 4
US-10-085-117-235/c
; Sequence 235, Application US/10085117
; Publication No. US20030232334A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Engelhard, Eric K.
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER
;  FILE REFERENCE: 529452000121
;  CURRENT APPLICATION NUMBER: US/10/085,117
;  CURRENT FILING DATE:  2002-02-27
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 361
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 235
;   LENGTH: 33488
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: variation
;   LOCATION: (1)...(33488)
;   OTHER INFORMATION: n = any nucleotide
US-10-085-117-235

  Query Match             1.8%;  Score 165;  DB 17;  Length 33488;
  Best Local Similarity   81.5%;  Pred. No. 6.6e-34;
  Matches  216;  Conservative    0;  Mismatches   45;  Indels    4;  Gaps    2;

Qy       4497 GCACTTGGGAGGCTGAAGCAGGAGGATTGCTATATGTTTGAGGCCAGCCTGAGCTATAGA 4556
              || || | ||||||||||||||| ||    ||  ||   |||||| |||| ||||   |
Db       6269 GCTCTGGTGAGGCTGAAGCAGGAGAAT---CATGAGTCCCAGGCCACCCTGGGCTACTTA 6213

Qy       4557 GCGAGACTTTGTCTTTAAGAAAAAAATG-AAAGCCCAGCAGTGGTGGCACACGCCTTTAA 4615
              | ||||| | ||||  || |||||||   |||||   || ||||| ||||||  |||||||
Db       6212 GTGAGACCTGGTCTAAAAAAAAAAAAAAAAAAAAGCTGGGCTGTGGTAGCACACACCTTTAA 6153

Qy       4616 TCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTAT 4675
              |||||||||||  ||| |||||  ||||||  ||||||||||||  ||||||||||||||||
Db       6152 TCCCAGCACTTTGGAAGCAGAGGCAGGCGGATTTCTGAGTTTGAGGCCAGCCTGGTCTAC 6093

Qy       4676 AGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAA 4735
              |||||||||||||||||||||||||||||||||||||||||||||| | ||||||  || |||
Db       6092 AGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTCTCGAAAAAACAAAAA 6033

Qy       4736 AACAAAACAAAACAAAACAAAACAA 4760
              || ||||||||  |||| |||| ||
Db       6032 AAAAAAACAAACAAAAAAAAAAAAA 6008


RESULT 5
US-10-034-650-1/c
; Sequence 1, Application US/10034650
; Publication No. US20030216558A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David
;  APPLICANT:  Engelhard, Eric
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
;  TITLE OF INVENTION:  CANCER
;  FILE REFERENCE: 529452000128
;  CURRENT APPLICATION NUMBER: US/10/034,650
;  CURRENT FILING DATE:  2002-07-23
;  PRIOR APPLICATION NUMBER: US 09/474,377
;  PRIOR FILING DATE: 1999-12-29
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 61
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1
;   LENGTH: 31632
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-10-034-650-1

  Query Match             1.8%;  Score 164;  DB 17;  Length 31632;
  Best Local Similarity   77.4%;  Pred. No. 1.2e-33;
  Matches  212;  Conservative    0;  Mismatches   60;  Indels    2;  Gaps    1;

Qy       4485 TGTATAATCTGAGCACTTGGGAGGCTGAAGCAGGAGGATTGCTATATGTTTGAGGCCAGC 4544
              ||  ||||| | || |  | | |||| |||| || |||| ||  |||| ||||||||
Db      20513 TGGGAAATCTCAACATTCAGAAAGCTGGGGCAGGTGGGTTGCCATGAGTTTCAGGCCAGC 20454

Qy       4545 CTGAGCTATAGAGCGAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCA 4604
              |||  |||||  |||||  ||| || || |  | ||    |||| ||||||||||||
Db      20453 CTGGCGTATAGTTTGAGACCCTGTTTTAAAAATGACAA--CCAGCCGGGCAGTGGTGGCA 20396

Qy       4605 CACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCA 4664
              || |||||||||||||||| |||||||||| |||||  |||||||||||| ||||||
Db      20395 TACACCTTTAATCCCAGCACTCGGGAGGCAGAGGCAGGAGGATTTCTGAGTTCGAGGCCA 20336

Qy       4665 GCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGA 4724
              |||||||||| | ||||||||||||||||||||||||| |||||||||| |||| |  |
Db      20335 GCCTGGTCTACAAAGTGAGTTCCAGGACAGCCAGGGCTATACAGAGAAACTCTGTCTCTA 20276

Qy       4725 AAAACCAGAAAAACAAAACAAAACAAAACAAAAC 4758
              |||||   || ||  ||| |||| |||| ||| |
Db      20275 AAAACTGAAAGAAGGAAAAAAAAAAAAAAAACC 20242


RESULT 6
US-10-322-696-25
; Sequence 25, Application US/10322696
; Publication No. US20040166490A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Malandro, Marc
;  TITLE OF INVENTION: NOVEL THERAPEUTIC TARGETS IN CANCER
;  FILE REFERENCE: 529452001200
;  CURRENT APPLICATION NUMBER: US/10/322,696
;  CURRENT FILING DATE:  2003-10-17
;  NUMBER OF SEQ ID NOS: 186
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 25
;   LENGTH: 21781
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-10-322-696-25

  Query Match             1.8%;  Score 161.4;  DB 19;  Length 21781;
  Best Local Similarity   81.0%;  Pred. No. 5.2e-33;
  Matches  200;  Conservative    0;  Mismatches   46;  Indels    1;  Gaps    1;

Qy       4516 AGGAGGATTGCTATATGTTTGAGGCCAGCCTGAGCTATAGAGCGAGACTTTGTCTTTAAG 4575
              ||  |||| || |  |||  ||  ||||| |||||| | || |||||  |||| | |||
Db       4520 AGATGGATCTCTGTGAGTTCAAGCACAGCCAGAGCTACACAGTGAGACCCTGTC-TCAAG 4578
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Qy       4576 AAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAG 4635
              ||   ||  ||||||  |  |||||||| ||||||||||||||||||||||||||||||||
Db       4579 AAGCGAACAAAAGCCGGGGCGTGGTGGCGCACGCCTTTAATCCCAGCACTTGGGAGGCAG 4638

Qy       4636 AAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGC 4695
              | |||||| |||||||||||||||||||||||||||||| | ||||||||||||||||||
Db       4639 AGGCAGGCGGATTTCTGAGTTCAAGGCCAGCCTGGTCTACATAGTGAGTTCCAGGACAGC 4698

Qy       4696 CAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAA 4755
              |||||||| ||||||||||||||| |  ||||| || | ||||||| || | |||| ||
Db       4699 CAGGGCTATACAGAGAAACCCTGTCTCAAAAAAACAAACAAACAAACAAAAAAAAA 4758

Qy       4756 AACAAAA 4762
                 |||||
Db       4759 CCCAAAA 4765
```

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RESULT 7
US-10-085-117-289
; Sequence 289, Application US/10085117
; Publication No. US20030232334A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Engelhard, Eric K.
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER
;  FILE REFERENCE: 529452000121
;  CURRENT APPLICATION NUMBER: US/10/085,117
;  CURRENT FILING DATE:  2002-02-27
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 361
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 289
;   LENGTH: 54648
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: variation
;   LOCATION: (1)...(54648)
;   OTHER INFORMATION: n = any nucleotide
US-10-085-117-289

  Query Match             1.8%;  Score 161.4;  DB 17;  Length 54648;
  Best Local Similarity   79.7%;  Pred. No. 9.2e-33;
  Matches  216;  Conservative    0;  Mismatches   51;  Indels    4;  Gaps    2;

Qy       4500 CTTGGGAGGCTGAAGCAGGAGGATTGCTATATGTTTGAGGCCAGCCTGAGCTATAGAGCG 4559
              | || || |   ||| | || || |||  |  |||||||||||| |||  ||||| |
Db      20809 CCTGAGAAGGATAAGGAAGAAGAATGCATTGAGTTTGAGGCCAGGCTGGACTATATAAAA 20868

Qy       4560 AGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCC 4619
              |     |   || || |||| |   |||||  |||||||||||| ||||||||||||||||
Db      20869 ACTAAGTAATTTAAACAAAACA---AAAGCGGGGCAGTGGTGGCGCACGCCTTTAATCCC 20925

Qy       4620 AGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAG 4679
              |||||||||||||||| |||||| |||||||||||||||||||||||||||||| ||||
Db      20926 AGCACTTGGGAGGCAGAGGCAGGCGGATTTCTGAGTTCAAGGCCAGCCTGGTCTACAGAG 20985

Qy       4680 TGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGA-AAAAC 4738
              |||||||||||||||||||||||||||||||||||||| | || |||||| |  ||||
Db      20986 TGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTCTCGATAAACCAAACCAAAC 21045

Qy       4739 AAAACAAAACAAAACAAAACAAAACCCAAAC 4769
              |||| |||| |||| |||| ||||| || |
Db      21046 CAAACCAAACCAAACCAAACCAAACCAAACC 21076
```

```
RESULT 8
US-10-322-281-51/c
; Sequence 51, Application US/10322281
; Publication No. US20040126762A1
; GENERAL INFORMATION:
;  APPLICANT: David W. Morris
;  APPLICANT: Marc S. Malandro
;  TITLE OF INVENTION: Novel Compositions and Methods in Cancer
;  FILE REFERENCE: 529452001000
;  CURRENT APPLICATION NUMBER: US/10/322,281
;  CURRENT FILING DATE:  2002-12-17
;  NUMBER OF SEQ ID NOS: 866
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 51
;   LENGTH: 111836
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(111836)
;   OTHER INFORMATION: n = A,T,C or G
US-10-322-281-51

  Query Match             1.8%;  Score 161;  DB 19;  Length 111836;
  Best Local Similarity   66.7%;  Pred. No. 1.9e-32;
  Matches  321;  Conservative    0;  Mismatches  100;  Indels   60;  Gaps    4;

Qy       1857 GGGAGAGATGGCTCAGCTTCCAGGAGCACTTGCTGCTCTTGCAGAGGACCTAGATTCAGT 1916
              ||||      | |||||      |  |||||| | |||||||  | ||||||||    ||  ||
Db      56644 GGGACTAGGGTCTCAGGGGTTAAAAGCACTGGTTGCTCTTCCTGAGGACCTGAGTTTGGT 56585

Qy       1917 TCCCAGGACTCATATGGTGGCTCACAGCCATCTGTAAATCCAGTTCCAGAGGGTTCCACA 1976
              ||||||  || ||||  ||  ||||||||  |  |||  |||  ||||||||   |  | | | |||
Db      56584 TCCCAGCACCCATACGGGGGCTCACAACTGTCTATAACTCCAGTTCTTGGAGATCCAACA 56525

Qy       1977 CCCTCTTCTGGCCTCCACAGGCACCACATACATAGTACACAGACATACATGCAGGCAAAA 2036
              |||||||||| ||||||  | ||  ||       | ||   |||| |||||  ||||||  |||
Db      56524 CCCTCTTCTGACCTCCATGGACATCAGGAGTGTGGTGTGCAGATATACAAGCAGGC-AAA 56466

Qy       2037 CACCCATACACACATAAATAAATAAGGAAACTTAAAAGGTGCATGTGTTGGTAAACATTG 2096
              |||  ||||||||    ||  ||| ||    |      |||||
Db      56465 CACTCATACACATCAAATAAAACAAATCAGAAAGAAAGG---------------------- 56427

Qy       2097 TGCTTACACATGCTGATTGAAGACATGTACAACGCACACACTGAAGAGGGATCTGGGGCT 2156
                                                  || | |  |  | ||       ||||||
Db      56426 --------------------------AAGGAAGGAAGGAAGGAAGGAAAAGAAGGGGGCT 56393

Qy       2157 GGAGAGATGGCTCAGCGGTTAAGAGCACTGACTGCTCTTCCGAAGGAAGGTCCTGAGTTC 2216
              |||||||||||||||||| ||||||||||||||||||||||     ||||||||||||||
Db      56392 GGAGAGATGGCTCAGCGGGTAAGAGCACTGACTGCTCTTCC----AAAGGTCCTGAGTTC 56337

Qy       2217 AAATCCTAGCAACCACATGGTGGCTCACAACCATCCATAATGAGATCTGACACCCTCTTC 2276
              ||||   |||||||||| |||||||||||||||| ||||||||||||||||||||||||||
Db      56336 AAATTACAGCAACCACATAGTGGCTCACAACCATTCATAATGAGATCTGACACCCTCTTC 56277

Qy       2277 TGGTGCATCTGAAGACAGCTGCAGAGCTACAGTGTACTTAGATATACTAATAAATAAATC 2336
              |||||||||||||| |        |||||||||||||||||| |||| ||||||||||||
Db      56276 TGGTGCATCTGAAGTC--------AGCTACAGTGTACTTAGGTATAATAATAAATAAATC 56225

Qy       2337 T 2337
              |
Db      56224 T 56224
```

```
RESULT 9
US-10-322-696-133
; Sequence 133, Application US/10322696
; Publication No. US20040166490A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Malandro, Marc
;  TITLE OF INVENTION: NOVEL THERAPEUTIC TARGETS IN CANCER
;  FILE REFERENCE: 529452001200
;  CURRENT APPLICATION NUMBER: US/10/322,696
;  CURRENT FILING DATE:  2003-10-17
;  NUMBER OF SEQ ID NOS: 186
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 133
;   LENGTH: 27383
```

```
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(27383)
;   OTHER INFORMATION: n = A,T,C or G
US-10-322-696-133

  Query Match             1.8%;  Score 158.8;  DB 19;  Length 27383;
  Best Local Similarity   81.4%;  Pred. No. 3.2e-32;
  Matches  184;  Conservative    0;  Mismatches   42;  Indels    0;  Gaps    0;

Qy       4559 GAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCC 4618
              || ||  | |||| || || |||   |    ||||  |||||||| ||||||||||||||
Db       6619 GATACCCTATCTTAAAAAACAAACAAACGAGCCAGGCGTGGTGGCGCACGCCTTTAATCC 6678

Qy       4619 CAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGA 4678
              ||||||| |||||||||| |||||| |||||||||||| |||||||||||||||| | |
Db       6679 CAGCACTCGGGAGGCAGAGGCAGGCGGATTTCTGAGTTCCAGGCCAGCCTGGTCTACAAA 6738

Qy       4679 GTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAAC 4738
              |||||||||||||||||||||||||||||||||||||||| ||||||||| | ||||||
Db       6739 GTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTCTTGAAAAACAAAAAAAAC 6798

Qy       4739 AAAACAAAACAAAACAAAACAAAACCCAAACCCAAACCCAAACCTC 4784
              ||||  |||| ||| |||  || |  |||||  | |  |||| | |
Db       6799 AAAAACAAACCAAAAAAACAAACAAACAAACAAACAAAACCC 6844


RESULT 10
US-10-087-192-1705
; Sequence 1705, Application US/10087192
; Publication No. US20020182586A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Engelhard, Eric K.
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
;  TITLE OF INVENTION:  CANCER
;  FILE REFERENCE: 529452000122
;  CURRENT APPLICATION NUMBER: US/10/087,192
;  CURRENT FILING DATE:  2002-03-01
;  PRIOR APPLICATION NUMBER: US 09/747,377
;  PRIOR FILING DATE: 2000-12-22
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 2059
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1705
;   LENGTH: 26345
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-10-087-192-1705

  Query Match             1.8%;  Score 158;  DB 13;  Length 26345;
  Best Local Similarity   80.4%;  Pred. No. 5.3e-32;
  Matches  185;  Conservative    0;  Mismatches   45;  Indels    0;  Gaps    0;

Qy       4532 GTTTGAGGCCAGCCTGAGCTATAGAGCGAGACTTTGTCTTTAAGAAAAAAATGAAAGCCC 4591
              |||  | |  |||  |  ||| | || || ||  |||||||| || ||||  |    ||
Db      18508 GTTCCACGATAGCTAGGACTACACAGAGAAACCCTGTCTTTAAAAACAAAACAAGGAGCC 18567

Qy       4592 AGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCT 4651
              ||  |||||||| |||||||||||||| |||||||||||||||| |||||| | |||||
Db      18568 AGTCGTGGTGGCGCACGCCTTTAATCCTAGCACTTGGGAGGCAGAGGCAGGCGGGTTTCT 18627

Qy       4652 GAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGA 4711
              |||||| |||||||||||||| |||||||||||||||||||||||||||||||||||||
Db      18628 GAGTTCGAGGCCAGCCTGGTCTACAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGA 18687

Qy       4712 AACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAA 4761
              |||||||| | |||||||| |||||  |||||    | ||| | ||| | |
Db      18688 AACCCTGTCTCGAAAAACCTTAAAAAAAAAACCCCAAAAATCCAAAAAGA 18737


RESULT 11
US-10-087-192-1663/c
; Sequence 1663, Application US/10087192
; Publication No. US20020182586A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Engelhard, Eric K.
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
;  TITLE OF INVENTION:  CANCER
;  FILE REFERENCE: 529452000122
;  CURRENT APPLICATION NUMBER: US/10/087,192
;  CURRENT FILING DATE:  2002-03-01
;  PRIOR APPLICATION NUMBER: US 09/747,377
;  PRIOR FILING DATE: 2000-12-22
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 2059
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1663
;   LENGTH: 193853
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(193853)
;   OTHER INFORMATION: n = A,T,C or G
US-10-087-192-1663

  Query Match             1.8%;  Score 157.6;  DB 13;  Length 193853;
  Best Local Similarity   85.8%;  Pred. No. 2.4e-31;
  Matches  175;  Conservative    0;  Mismatches   29;  Indels    0;  Gaps    0;

Qy       4559 GAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCC 4618
              ||||  ||   ||||| |   ||     ||||  ||||||||||||||||||||||||||
Db     123834 GAGAACTTTGTTTTAAAAGGCAACAATGAGCCAGGCAGTGGTGGCACACGCCTTTAATCC 123775

Qy       4619 CAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGA 4678
              |||||||||||||||| |||||| |||||||||||||||||||||||||||||| |||
Db     123774 TAGCACTTGGGAGGCAGAGGCAGGCGGATTTCTGAGTTCAAGGCCAGCCTGGTCTACAGA 123715

Qy       4679 GTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAAC 4738
              |||||||||||||||| ||||||||||||||||||||||| | ||||||||| |||||
Db     123714 GTGAGTTCCAGGACAGTCAGGGCTACACAGAGAAACCCTGTCTCGAAAAACCAAAAAAAA 123655

Qy       4739 AAAACAAAACAAAACAAAACAAAA 4762
              |||| |||| |||| |||| ||||
Db     123654 AAAAAAAAAAAAAAAAAAAAAAAA 123631


RESULT 12
US-10-322-281-817/c
; Sequence 817, Application US/10322281
; Publication No. US20040126762A1
; GENERAL INFORMATION:
;  APPLICANT: David W. Morris
;  APPLICANT:  Marc S. Malandro
;  TITLE OF INVENTION: Novel Compositions and Methods in Cancer
;  FILE REFERENCE: 529452001000
;  CURRENT APPLICATION NUMBER: US/10/322,281
;  CURRENT FILING DATE:  2002-12-17
;  NUMBER OF SEQ ID NOS: 866
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 817
;   LENGTH: 93329
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(93329)
;   OTHER INFORMATION: n = A,T,C or G
US-10-322-281-817
```

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  Query Match             1.7%;  Score 156.2;  DB 19;  Length 93329;
  Best Local Similarity   78.0%;  Pred. No. 3.8e-31;
  Matches  188;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy       4547 GAGCTATAGAGCGAGACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACA 4606
             ||  |||| | | | |   | | |   |  | ||||  ||||||  |  |||||||| ||
Db      32422 GAAATATATAACAAAATAATTTATAATAAGATAAAACAAAAGCCGGGTGGTGGTGGCGCA 32363

Qy       4607 CGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGC 4666
             ||||||||||||||||||||||||||||||||||  ||||||||||||| ||||||||
Db      32362 CGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGTGGATTTCTGAGTTCGAGGCCAGC 32303

Qy       4667 CTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAA 4726
             ||||||||  | |||||||||||||||||||||||||||| |||||||||||||| | ||||
Db      32302 CTGGTCTACAAAGTGAGTTCCAGGACAGCCAGGGCTATACAGAGAAACCCTGTCTCGAAA 32243

Qy       4727 AACCAGAAAAACAAAACAAAACAAAACAAAACAAAACCCAAACCCAAACCCAAACCTCTC 4786
             |||||  |||||  ||||  |   |   ||   | ||  ||||    ||     |||  | ||||||  ||
Db      32242 AACCAAAAAAAAAAAAAAACCAACAACAACAAAAAAAAAAAAACGATAAAACAAAACCTATC 32183

Qy       4787 A 4787
              |
Db      32182 A 32182


RESULT 13
US-10-367-094-142
; Sequence 142, Application US/10367094
; Publication No. US20040170982A1
; GENERAL INFORMATION:
;  APPLICANT: David W. Morris
;  APPLICANT:  Marc Malandro
;  TITLE OF INVENTION: Novel Therapeutic Targets in Cancer
;  FILE REFERENCE: 529452001500
;  CURRENT APPLICATION NUMBER: US/10/367,094
;  CURRENT FILING DATE:  2003-02-14
;  NUMBER OF SEQ ID NOS: 203
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 142
;   LENGTH: 24495
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(24495)
;   OTHER INFORMATION: n = A,T,C or G
US-10-367-094-142

  Query Match             1.7%;  Score 156;  DB 19;  Length 24495;
  Best Local Similarity   81.8%;  Pred. No. 1.9e-31;
  Matches  180;  Conservative    0;  Mismatches   40;  Indels    0;  Gaps    0;

Qy       4563 CTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGC 4622
             |||  | |  || |||||||  |||  || |  ||| || ||||||||||||||||||||
Db       2471 CTTGATTTAAAAAAAAAAAAAAAAAAAGCCGGGCGTGATGACACACGCCTTTAATCCCAGC 2530

Qy       4623 ACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGA 4682
             |||||||||||||| |||||  |||||||||||||||||||||||||| | |||||
Db       2531 ACTTGGGAGGCAGAGGCAGGTGGATTTCTGAGTTCAAGGCCAGCCTGGTCTACAAAGTGA 2590

Qy       4683 GTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAA 4742
             |||||||||||||||||||||||||||||||||||| |  |||||| | ||| | ||||
Db       2591 GTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTCTCCAAAAACAAAAAACAAAAAA 2650

Qy       4743 CAAAACAAAACAAAACAAAACCCAAACCCAAACCCAAACC 4782
             | |||| |||| |||| ||||| | | | | | | | | |
Db       2651 CCAAACCAAACCAAACCAAACCAACAACAACAACAAGAAC 2690


RESULT 14
US-10-087-192-907
; Sequence 907, Application US/10087192
; Publication No. US20020182586A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David W.
;  APPLICANT:  Engelhard, Eric K.
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
;  TITLE OF INVENTION:  CANCER
;  FILE REFERENCE: 529452000122
;  CURRENT APPLICATION NUMBER: US/10/087,192
;  CURRENT FILING DATE:  2002-03-01
;  PRIOR APPLICATION NUMBER: US 09/747,377
;  PRIOR FILING DATE: 2000-12-22
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 2059
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 907
;   LENGTH: 31400
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(31400)
;   OTHER INFORMATION: n = A,T,C or G
US-10-087-192-907

  Query Match             1.7%;  Score 155.6;  DB 13;  Length 31400;
  Best Local Similarity   87.6%;  Pred. No. 2.8e-31;
  Matches  170;  Conservative    0;  Mismatches   24;  Indels    0;  Gaps    0;

Qy       4587 AGCCCAGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAAGCAGGCAGA 4646
             ||||  |||||||||||||||||||||||||||||||||||||||||||||||| ||||| |||
Db        294 AGCCAGGCAGTGGTGGCACACGCCTTTAATCCCAGCACTTGGGAGGCAGAGGCAGGTAGA 353

Qy       4647 TTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGTGAGTTCCAGGACAGCCAGGGCTACAC 4706
             |||||||||||| |||||||||||||||| |||||||||||||||||||||||||||||||
Db        354 TTTCTGAGTTCGAGGCCAGCCTGGTCTACAGAGTGAGTTCCAGGACAGCCAGGGCTACAC 413

Qy       4707 AGAGAAACCCTGTTTTGAAAAACCAGAAAAACAAAACAAAACAAAACAAAACAAAACCCA 4766
             |||||||||||||| |||||||||| ||||| |||| |||| |||| |||| |||| |||
Db        414 AGAGAAACCCTGTCTTGAAAAACCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAGAGTT 473

Qy       4767 AACCCAAACCCAAA 4780
              |  || ||  |||
Db        474 GAATCATACATAAA 487


RESULT 15
US-09-997-722-31
; Sequence 31, Application US/09997722
; Publication No. US20040072154A1
; GENERAL INFORMATION:
;  APPLICANT: Morris, David
;  APPLICANT:  Engelhard, Eric
;  TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER
;  FILE REFERENCE: A-71171/RMS/DCF
;  CURRENT APPLICATION NUMBER: US/09/997,722
;  CURRENT FILING DATE:  2001-11-30
;  PRIOR APPLICATION NUMBER: US 09/747,377
;  PRIOR FILING DATE: 2000-12-22
;  PRIOR APPLICATION NUMBER: US 09/798,586
;  PRIOR FILING DATE: 2001-03-02
;  NUMBER OF SEQ ID NOS: 301
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31
;   LENGTH: 94381
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (2681)..(2700)
;   OTHER INFORMATION: "n" at positions 2681 through 2700 can be any base.
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (11224)..(11290)
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; OTHER INFORMATION: "n" at positions 11224 through 11290 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (28032)..(28051)
; OTHER INFORMATION: "n" at positions 28032 through 28051 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (37054)..(37178)
; OTHER INFORMATION: "n" at positions 37054 through 37178 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (44115)..(45641)
; OTHER INFORMATION: "n" at positions 44115 through 45641 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (47999)..(48255)
; OTHER INFORMATION: "n" at positions 47999 through 48255 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (59012)..(62394)
; OTHER INFORMATION: "n" at positions 59012 through 62394 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (63574)..(63720)
; OTHER INFORMATION: "n" at positions 63574 through 63720 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (70645)..(70664)
; OTHER INFORMATION: "n" at positions 70645 through 70664 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (85078)..(85097)
; OTHER INFORMATION: "n" at positions 85078 through 85097 can be any base.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (93428)..(93447)
; OTHER INFORMATION: "n" at positions 93428 through 93447 can be any base.

US-09-997-722-31

Query Match 1.7%; Score 155; DB 11; Length 94381;
Best Local Similarity 80.2%; Pred. No. 8.3e-31;
Matches 182; Conservative 0; Mismatches 45; Indels 0; Gaps 0;

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Qy       4561 GACTTTGTCTTTAAGAAAAAAATGAAAGCCCAGCAGTGGTGGCACACGCCTTTAATCCCA 4620
              |||  |  ||||||  ||  | | |||    ||||  |||||||| ||||||||||||||||
Db      30936 GACAGTCCCTTTAAAAACACAGTGACTAGCCAGGTGTGGTGGCGCACGCCTTTAATCCCA 30995

Qy       4621 GCACTTGGGAGGCAGAAGCAGGCAGATTTCTGAGTTCAAGGCCAGCCTGGTCTATAGAGT 4680
              ||||||||||||||| ||||||||||||||||||  ||||||||||||||| | |||
Db      30996 GCACTTGGGAGGCAGAGGCAGGCAGATTTCTGAGTTTGAGGCCAGCCTGGTCTACAAAGT 31055

Qy       4681 GAGTTCCAGGACAGCCAGGGCTACACAGAGAAACCCTGTTTTGAAAAACCAGAAAAACAA 4740
              |||||||||||||||||||||| ||||||||||||||| | ||| ||||| ||||| ||
Db      31056 GAGTTCCAGGACAGCCAGGGCTATACAGAGAAACCCTGTCTCGAATAACCAAAAAAAAAA 31115

Qy       4741 AACAAAACAAAACAAAACAAAACCCAAACCCAAACCCAAACCTCTCA 4787
              || |||| ||| ||||| || |  | || |  |  |  |  | | ||
Db      31116 AAAAAAAAAAACCAAAAAAACAAACCAAACACACACACACACACA 31162
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Search completed: October 18, 2005, 13:24:01
Job time : 5404 secs

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